	umber: 09186662	CRF Processing Edited by:		
1 13	Changed a file from non-ASCII to ASCII	Verified by:	DC	(STIC st
	Changed the margins in cases where the sequence text was "wrapping to the County Application Data socion, specifical			n 1
	Edited a format error in the Current Application Data section, specifical	lly:		
	Edited the Current Application Data section with the actual current numapplicant was the prior application data; or other	nber. The numb	er inputte	ed by the
	Added the mandatory heading and subheadings for *Current Applicatio	n Data*.		
	Edited the "Number of Sequences" field. The applicant spelled out a nu	umber instead c	If using a	n integer.
	Changed the spelling of a mandatory field (the headings or subheading	s), specifically:	-	
	Corrected the SEQ ID NO when obviously incorrect. The sequence number of the sequence of the s	mbers that were	edited w	rere:
	Inserted or corrected a nucleic number at the end of a nucleic line. SE	Q ID NO's edite	d:	
	Corrected subheading placement. All responses must be on the same lapplicant placed a response below the subheading, this was moved to i	line as each sut ts appropriate p	heading. lace.	If the
	Inserted colons after headings/subheadings. Headings edited included			
	Deleted extra, invalid, headings used by an applicant, specifically:			
	Deleted: non-ASCII "garbage" at the beginning/end of files; se page numbers throughout text; other invalid text, such as	ecretary initials/fi	lename a	at end of file
	Inserted mandatory headings, specifically:		· · · · · · · · · · · · · · · · · · ·	
	Corrected an obvious error in the response, specifically:			
	Edited identifiers where upper case is used but lower case is required.	or vice versa.		
	Corrected an error in the Number of Sequences field, specifically:			۰
_	A "Hard Page Break" code was inserted by the applicant. All occurrence	ces had to be do	leted.	
	Peleted ending stop codon in amino acid sequences and adjusted the function to a Patentin bug). Sequences corrected:			ngly (error
	Other: For Say 90+91 moved agail 4 ago-12, respectively	y pa 22237,	to previ	· y line

Petch

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

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RAW SEQUENCE LISTING DATE: 11/07/2001 PATENT APPLICATION: US/09/856,662 TIME: 09:18:51

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\11072001\1856662.raw

- 3 <110> APPLICANT: MORIBE, Toyoki et al.
  5 <120> TITLE OF INVENTION: Method for typing HLA class 1 genes
- 7 <130> FILE REFERENCE: 0032-0261P
- 9 <140> CURRENT APPLICATION NUMBER: US 09/856,662
- 10 <141> CURRENT FILING DATE: 2001-05-24
- 12 <150> PRIOR APPLICATION NUMBER: JP P1998-335151
- 13 <151> PRIOR FILING DATE: 1998-11-26
- 15 <160> NUMBER OF SEQ ID NOS: 130
- 17 <170> SOFTWARE: PatentIn Ver. 2.0
- 19 <210> SEQ ID NO: 1
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- 28 gaggtatttc ttcacatccg tgt
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- 54 <212> TYPE: DNA
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- 65 <212> TYPE: DNA
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RAW SEQUENCE LISTING

DATE: 11/07/2001

PATENT APPLICATION: US/09/856,662

TIME: 09:18:51

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- 71 <400> SEQUENCE: 5
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- 74 <210> SEQ ID NO: 6
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- 131 <210> SEQ ID NO: 11 132 <211> LENGTH: 20
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RAW SEQUENCE LISTING DATE: 11/07/2001 PATENT APPLICATION: US/09/856,662 TIME: 09:18:51

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Output Set: N:\CRF3\11072001\1856662.raw

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RAW SEQUENCE LISTING

DATE: 11/07/2001 TIME: 09:18:51

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PATENT APPLICATION: US/09/856,662

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\11072001\1856662.raw

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- 205 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA probe A368A
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- 210 <210> SEQ ID NO: 18
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- 212 <212> TYPE: DNA
- 213 <213> ORGANISM: Artificial Sequence
- 215 <220> FEATURE:
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- 230 gaggatgttt ggctgc
- 232 <210> SEQ ID NO: 20
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- 234 <212> TYPE: DNA
- 235 <213> ORGANISM: Artificial Sequence
- 237 <220> FEATURE:
- 238 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA probe A402G
- 240 <400> SEQUENCE: 20
- 241 cgcttcctgc gcgggt
- 243 <210> SEQ ID NO: 21
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- 252 caggacgctt acgacgg
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- 265 <210> SEQ ID NO: 23
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- 267 <212> TYPE: DNA
- 268 <213> ORGANISM: Artificial Sequence
- 270 <220> FEATURE:

RAW SEQUENCE LISTING DATE: 11/07/2001 PATENT APPLICATION: US/09/856,662 TIME: 09:18:51

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\11072001\1856662.raw

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337 <220> FEATURE:

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/856,662

DATE: 11/07/2001

TIME: 09:18:52

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\11072001\1856662.raw

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RAW SEQUENCE LISTING
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                     PATENT APPLICATION: US/09/856,662
                                                              TIME: 13:07:49
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      5 <120 > TITLE OF INVENTION: Method for typing HLA class 1 genes
      7 <130 FILE REFERENCE: 0032-0261P
      9 <140 > CURRENT APPLICATION NUMBER: US 09/856,662
     10 <141> CURRENT FILING DATE: 2001-05-24
     12 <150> PRIOR APPLICATION NUMBER: JP P1998-335151
     13 <151> PRIOR FILING DATE: 1998-11-26
                                                                                        ور الإلكار ال
     15 <160> NUMBER OF SEO ID NOS: 130
     17 <170> SOFTWARE: PatentIn Ver. 2.0
                                                             Cestion compris
ERRORED SEQUENCES
                                                             Careano Chakette Nesage
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     1024 <211> LENGTH: 19
     1025 - 1212> TYPE: DNA
     1026 <213> ORGANISM: Artificial Sequence
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E--> 1030
                               incred each it is to fine above to good computer mi interpretation
W--> 1032 <210> SEQ ID NO:
W--> 1032 <211> LENGTH:
W--> 1032 <212> TYPE:
W--> 1032 <213> ORGANISM:
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## VERIFICATION SUMMARY

DATE: 10/29/2001 TIME: 13:07:51 PATENT APPLICATION: US/09/856,662

Input Set : A:\0032-0261P.ST25.txt

Output Set: N:\CRF3\10292001\1856662.raw

L:1030 M:254 E: No. of Bases conflict, LENGTH:Input:11 Counted:3 SEQ:90 L:1032 M:282 W: Numeric Field Identifier Missing, <210> is required. L:1032 M:282 W: Numeric Field Identifier Missing, <211> is required. L:1032 M:282 W: Numeric Field Identifier Missing, <212> is required. L:1032 M:282 W: Numeric Field Identifier Missing, <213> is required. L:1042 M:254 E: No. of Bases conflict, LENGTH:Input:12 Counted:3 SEQ:91 M:112 Repeated in SeqNo=91 L:1042 M:352 E: No. of Seq. differs, <211>LENGTH:Input:19 Found:3 SEQ:91 L:1044 M:282 W: Numeric Field Identifier Missing, <210> is required. L:1044 M:282 W: Numeric Field Identifier Missing, <211> is required. L:1044 M:282 W: Numeric Field Identifier Missing, <212> is required. L:1044 M:282 W: Numeric Field Identifier Missing, <213> is required.